PATENT

Docket No.: 176/60792 (6-11415-868)

Examiner: Unknown

Art Unit:

Unknown

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant

Mahin D. Maines

Serial No.

09/606,129

Filed

For

June 28, 2000

BILIVERDIN REDUCTASE FRAGMENTS

AND VARIANTS, AND METHODS OF USING BILIVERDIN REDUCTASE AND SUCH FRAGMENTS AND VARIANTS

SUBMISSION OF REVISED SEQUENCE LISTING AND STATEMENT IN ACCORDANCE WITH 37 C.F.R. § 1.821(g)

Assistant Commissioner for Patents Washington, D.C. 20231

Sir:

In response to the Notice to File Missing Parts, applicant hereby submits a revised Sequence Listing (21 pages) on paper and on a computer readable 3.5" diskette. The revised sequence listing differs from the originally filed sequence listing by including data for the present application and revising the description of SEQ. ID. Nos. 7 and 8 to indicate that X at each position can be any amino acid. For each of SEQ. ID. Nos. 6-17, the listed sequences are motifs which generally define the sequences of various hGVR domains; hence, an X at each position can be any amino acid. In the originally filed sequence listing, typographical errors resulted in the incomplete description of SEQ. ID. Nos. 7 and 8. The revised Sequence Listing corrects these errors.

Pursuant to 37 CFR § 1.821(g), applicant submits that the information on the paper copy and the computer readable diskette are the same. This submission contains no new matter.

Dated: Pecember 18, 2000

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R431370.1

Respectfully submitted,

Edwin V. Merkel

Registration No. 40,087

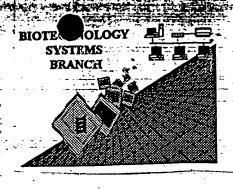
Certificate of Mailing - 37 CFR 1.8 (a)

I hereby certify that this correspondence is being deposited with the United States Postal Service as first class mail in an envelope addressed to: Assistant Commissioner for Patents, Washington, D.C. 20231. on the

Datě

Wendy L. Harrold

RAW-SEQUENCE-LISTENGI PERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/606,/29

Source: O/Pi

Date Processed by STIC: $\frac{7/12/2000}{}$

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR FURTHER INFORMATION, PLEASE TELEPHONE MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

OIPE

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DATE: 07/12/2000
RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/606,129
                                        TIME: 10:58:14
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Input Set : A:\U607921.app

Output Set: N:\CRF3\07122000\I606129.raw

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<120> TITLE OF INVENTION: BILIVERDIN REDUCTASE FRAGMENTS AND VARIANTS, AND METHODS OF USING RILIVERDIN PROJECTION AND COMMENTS.
       3 <110> APPLICANT: Maines, Mahin D.
               METHODS OF USING BILIVERDIN REDUCTASE AND SUCH
               FRAGMENTS AND VARIANTS
      9 <130> FILE REFERENCE: 176/60792
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/606,129
C--> 12 <141> CURRENT FILING DATE: 2000-06-28
     14 <150> PRIOR APPLICATION NUMBER: 60/141,309
     15 <151> PRIOR FILING DATE: 1999-06-28
                                                                  Does Not Comply
     17 <150> PRIOR APPLICATION NUMBER: 60/163,223
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     18 <151> PRIOR FILING DATE: 1999-11-03
     20 <160> NUMBER OF SEQ ID NOS: 37
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     24 <210> SEQ ID NO: 1
     25 <211> LENGTH: 296
     26 <212> TYPE: PRT
     27 <213> ORGANISM: Homo sapiens
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     36 Ser Ser Ala Phe Leu Asn Leu Ile Gly Phe Val Ser Arg Arg Glu Leu 37 \phantom{\bigg|}40\phantom{\bigg|} 40 \phantom{\bigg|}45\phantom{\bigg|}
     39 Gly Ser Ile Asp Gly Val Gln Gln Ile Ser Leu Glu Asp Ala Leu Ser 40 50 60
     42 Ser Gln Glu Val Glu Val Ala Tyr Ile Cys Ser Glu Ser Ser Ser His 43 65 70 75 80
     45 Glu Asp Tyr Ile Arg Gln Phe Leu Asn Ala Gly Lys His Val Leu Val 46 90 95
     48 Glu Tyr Pro Met Thr Leu Ser Leu Ala Ala Ala Glu Glu Leu Trp Glu 49 100 105 110
                                          105
     51 Leu Ala Glu Gln Lys Gly Lys Val Leu His Glu Glu His Val Glu Leu
                                      120
     54 Leu Met Glu Glu Phe Ala Phe Leu Lys Lys Glu Val Val Gly Lys Asp
                                  135
            130
     57 Leu Leu Lys Gly Ser Leu Leu Phe Thr Ser Asp Pro Leu Glu Glu Asp
                              150
                                                    155
     58 145
     60 Arg Phe Gly Phe Pro Ala Phe Ser Gly Ile Ser Arg Leu Thr Trp Leu
                         165
                                               170
     63 Val Ser Leu Phe Gly Glu Leu Ser Leu Val Ser Ala Thr Leu Glu Glu
                    180
                                         185
     66 Arg Lys Glu Asp Gln Tyr Met Lys Met Thr Val Cys Leu Glu Thr Glu
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200

21'5

69 Lys Lys Ser Pro Leu Ser Trp Ile Glu Glu Lys Gly Pro Gly Leu Lys

72 Arg Asn Arg Tyr Leu Ser Phe His Phe Lys Ser Gly Ser Leu Glu Asn

205

195

RAW SEQUENCE LISTING DATE: 07/12/2000 PATENT APPLICATION: US/09/606,129 TIME: 10:58:14

Input Set : A:\U607921.app

Output Set: N:\CRF3\07122000\1606129.raw

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73 225
                       230
                                           235
75 Val Pro Asn Val Gly Val Asn Lys Asn Ile Phe Leu Lys Asp Gln Asn
                 245
                                     250
78 Ile Phe Val Gln Lys Leu Leu Gly Gln Phe Ser Glu Lys Glu Leu Ala
                                  265
                                                      270
             260
81 Ala Glu Lys Lys Arg Ile Leu His Cys Leu Gly Leu Ala Glu Glu Ile
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84 Gln Lys Tyr Cys Cys Ser Arg Lys
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96 tccgtgcgga tgagggactt gcggaatcca caccetteet cagegtteet gaacetgatt 180
97 ggcttcgtgt cgagaaggga gctcgggagc attgatggag tccagcagat ttctttggag 240
98 gatgetettt eeageeaaga ggtggaggte geetatatet geagtgagag eteeageeat 300
99 gaggaetaca teaggeagtt eettaatget ggeaageaeg teettgtgga atacceeatg 360
100 acactgtcat tggcggccgc tcaggaactg tgggagctgg ctgagcagaa aggaaaagtc 420
101 ttgcacgagg agcatgttga actcttgatg gaggaattcg ctttcctgaa aaaagaagtg 480
102 gtggggaaag acctgctgaa agggtcgctc ctcttcacat ctgacccgtt ggaagaagac 540
103 eggtttgget teeetgeatt cageggeate tetegaetga eetggetggt etecetettt 600
104 ggggagettt etettgtgte tgecaetttg gaagagegaa aggaagatea gtatatgaaa 660
105 atgacagtgt gtctggagac agagaagaaa agtccactgt catggattga agaaaaagga 720
106 cctggtctaa aacgaaacag atatttaagc ttccatttca agtctgggtc cttggagaat 780
107 gtgccaaatg taggagtgaa taagaacata tttctgaaag atcaaaatat atttgtccag 840
108 aaactettgg gecagttete tgagaaggaa etggetgetg aaaagaaacg cateetgeae 900
109 tgcctggggc ttgcagaaga aatccagaaa tattgctgtt caaggaagta agaggaggag 960
110 gtgatgtagc acttccaaga tggcaccagc atttggttct tctcaagagt tgaccattat 1020
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115 <211> LENGTH: 296
116 <212> TYPE: PRT
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126 Ser Ser Ala Phe Leu Asn Leu Ile Gly Phe Val Ser Arg Arg Glu Leu
            3.5
                                40
129 Gly Ser Ile Asp Gly Val Gln Gln Ile Ser Leu Glu Asp Ala Leu Ser
130
      50
                             55
132 Ser Gln Glu Val Glu Val Ala Tyr Ile Cys Ser Glu Ser Ser Ser His
                                          75
                     70
135 Glu Asp Tyr Ile Arg Gln Phe Leu Asn Ala Gly Lys His Val Leu Val
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RAW SEQUENCE LISTING DATE: 07/12/2000 PATENT APPLICATION: US/09/606,129 TIME: .0:58:14

Input Set : A:\U607921.app

Output Set: N:\CRF3\07122000\1606129.raw

138 Glu Tyr Pro Met Thr Leu Ser Leu Ala Ala Ala Gln Glu Leu Trp Glu 100 105 141 Leu Ala Glu Gln Lys Gly Lys Val Leu His Glu Glu His Val Glu Leu 115 120 125 144 Leu Met Glu Glu Phe Ala Phe Leu Lys Lys Glu Val Val Gly Lys Asp 145 130 135 140 147 Leu Leu Lys Gly Ser Leu Leu Phe Thr Ala Gly Pro Leu Glu Glu Glu 148 145 150 150 155 160 150 Arg Phe Gly Phe Pro Ala Phe Ser Gly Ile Ser Arg Leu Thr Trp Leu 151 165 170 175 153 Val Ser Leu Phe Gly Glu Leu Ser Leu Val Ser Ala Thr Leu Glu Glu 154 180 185 190 180 156 Arg Lys Glu Asp Gln Tyr Met Lys Met Thr Val Cys Leu Glu Thr Glu 157 200 205 159 Lys Lys Ser Pro Leu Ser Trp Ile Glu Glu Lys Gly Pro Gly Leu Lys 160 210 215 220 162 Arg Asn Arg Tyr Leu Ser Phe His Phe Lys Ser Gly Ser Leu Glu Asn 163 225 230 235 240 165 Val Pro Asn Val Gly Val Asn Lys Asn Ile Phe Leu Lys Asp Gln Asn 166 245 250 255 168 Ile Phe Val Gln Lys Leu Leu Gly Gln Phe Ser Glu Lys Glu Leu Ala 169 265 270 171 Ala Glu Lys Lys Arg Ile Leu His Cys Leu Gly Leu Ala Glu Glu Ile 172 275 280 285 174 Gln Lys Tyr Cys Cys Ser Arg Lys 175 290 295 178 <210> SEQ ID NO: 4 179 <211> LENGTH: 295 180 <212> TYPE: PRT 181 <213> ORGANISM: Rattus norvegicus 183 <400> SEQUENCE: 4 184 Met Asp Ala Glu Pro Lys Arg Lys Phe Gly Val Val Val Val Gly Val
185 1 5 10 15 187 Gly Arg Ala Gly Ser Val Arg Leu Arg Asp Leu Lys Asp Pro Arg Ser 188 20 193 Ser Leu Asp Glu Val Arg Gln Ile Ser Leu Glu Asp Ala Leu Arg Ser 194 55 55 60 196 Gln Glu Ile Asp Val Ala Tyr Ile Cys Ser Glu Ser Ser Ser His Glu 197 65 70 75 80 199 Asp Tyr Ile Arg Gln Phe Leu Gln Ala Gly Lys His Val Leu Val Glu 200 85 90 95 202 Tyr Pro Met Thr Leu Ser Phe Ala Ala Ala Gln Glu Leu Trp Glu Leu 203 100 105 . 110 205 Ala Ala Gln Lys Gly Arg Val Leu His Glu Glu His Val Glu Leu Leu 115 120 125 208 Met Glu Glu Phe Glu Phe Leu Arg Arg Glu Val Leu Gly Lys Glu Leu

135

PATENT APPLICATION: US/09/606,129 TIME: 10:58:14 Input Set : A:\U607921.app Output Set: N:\CRF3\07122000\1606129.raw 211 Leu Lys Gly Ser Leu Arg Phe Thr Ala Ser Pro Leu Glu Glu Glu Arg 212 145 150 214 Phe Gly Phe Pro Ala Phe Ser Gly Ile Ser Arg Leu Thr Trp Leu Val 165 . 170 Ser Leu Phe Gly Glu Leu Ser Leu Ile Ser Ala Thr Leu Glu Glu Arg 180 185 190 218 220 Lys Glu Asp Gln Tyr Met Lys Met Thr Val Gln Leu Glu Thr Gln Asn 221 195 200 205 223 Lys Gly Leu Leu Ser Trp Ile Glu Glu Lys Gly Pro Gly Leu Lys Arg 210 215 220 224 226 Asn Arg Tyr Val Asn Phe Gln Phe Thr Ser Gly Ser Leu Glu Glu Val 227 225 230 235 240 229 Pro Ser Val Gly Val Asn Lys Asn Ile Phe Leu Lys Asp Gln Asp Ile 230 245 250 255245 250 232 Phe Val Gln Lys Leu Leu Asp Gln Val Ser Ala Glu Asp Leu Ala Ala 233 260 265 270 235 Glu Lys Lys Arg Ile Met His Cys Leu Gly Leu Ala Ser Asp Ile Gln 236 275 280 285 238 Lys Leu Cys His Gln Lys Lys 239 290 295 290 239 242 <210> SEQ ID NO: 5 243 <211> LENGTH: 1081 244 <212> TYPE: DNA 245 <213> ORGANISM: Rattus norvegicus 247 <400> SEQUENCE: 5 248 ggtcaacage taagtgaage catatecata gagagtttgt gecagtgeee caagateetg 60 249 aacctctgtc tgtcttcgga cactgactga agagaccgag atggatgccg agccaaagag 120 250 gaaatttgga gtggtagtgg ttggtgttgg cagagctggc tcggtgaggc tgagggactt 180 251 gaaggateca egetetgeag catteetgaa eetgattgga ttigtgteea gaegagaget 240 252 tgggagcctt gatgaagtac ggcagattte tttggaagat gcteteegaa gccaagagat 300 253 tgatgtegee tatattigea gtgagagtte eagecatgaa gaetatatae ggeagtttet 360 254 geaggetgge aageatgtee tegtggaata eeccatgaca etgteatttg eggeggeeca 420 255 ggagetgtgg gagetggeeg caeagaaagg gagagteetg catgaggage aegtggaaet 480 256 cttgatggag gaattcgaat tcctgagaag agaagtgttg gggaaagagc tactgaaagg 540 257 gtotottogo ticacagota goccactgga agaagagaa titiggottoo otgogitoag 600 258 eggeatttet egeetyacet ggetggtete eetetteggg gagetttete tratttetge 660 259 caccttggaa gagcgaaang aggatcagta tatgaaaatg accgtgcagc tggagaccca 720 260 gaacaagggt etgetgteat ggattgaaga gaaagggeet ggettaaaaa gaaacagata 780 261 tgtaaacttc cagttcactt ctgggtccct ggaggaagtg ccaagtgtag gggtcaataa 840 262 gaacattttc ctgaaagatc aggatatatt tgttcagaag ctcttagacc aggtctctgc 900 263 agaggacetg gefgetgagu agaagegeat cafgeattge etggggetgg ceagegacat 960 264 eeagaagett tgeeaceaga agaagtgaag aggaagette agagacettet gaagggggee 1020 265 agggtttggt cotatcaacc attcaccttt agctcttaca attaaacaty tcagataaac 1060 266 a 269 <210> SEQ ID NO: 6 270 <211> LENGTH: 6

DATE: 07/12/2000

RAW SEQUENCE LISTING

271 <212> TYPE: PRT

274 <220> FEATURE:

272 <213> ORGANISM: Artificial Sequence

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TIME: 10:58:14
                     PATENT APPLICATION: US/09/606,129
                     Input Set : A:\U607921.app
                     Output Set: N:\CRF3\07122000\1606129.raw
     275 <223> OTHER INFORMATION: Description of Artificial Sequence: hydrophobic
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     278 <220> FEATURE:
     279 <221> NAME/KEY: PEPTIDE
     280 <222> LOCATION: (2)
     281 <223> OTHER INFORMATION: where X is any aa
     283 <400> SEQUENCE: 6
284 Phe Xaa Val Val Val Val
     285
     288 <210> SEQ ID NO: 7
     289 <211> LENGTH: 6
     290 <212> TYPE: PRT
     291 <213> ORGANISM: Artificial Sequence
     293 <220> FEATURE:
     294 <223> OTHER INFORMATION: Description of Artificial Sequence: nucleotide
     295
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     297 <220> FEATURE:
     298 <221> NAME/KEY: PEPTIDE
     299 <222> LOCATION: (2)
     300 <223> OTHER INFORMATION: where X is any aa
     302 <400> SEQUENCE: 7
(W) -> 303 Gly Xaa Gly Xaa Xaa Gly
     304 1
     307 <210> SEQ ID NO: 8
     308 <211> LENGTH: 8
     309 <212> TYPE: PRT
     310 <213> ORGANISM: Artificial Sequence
     312 <220> FEATURE:
     313 <223> OTHER INFORMATION: Description of Artificial Sequence:
     314
               oxidoreductase domain of BVR
     316 <400> SEQUENCE: 8
     317 Ala Gly Leu His Val Leu Val Glu
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     321 <210> SEQ ID NO: 9
     322 <211> LENGTH: 29
     323 <212> TYPE: PRT
     324 <213> ORGANISM: Artificial Sequence
     326 <220> FEATURE:
     327 <223> OTHER INFORMATION: Description of Artificial Sequence: leucine
                                                etabout Xaa's at location 3
     328
               zipper of BVR
     330 <220> FEATURE:
     331 <221> NAME/KEY: PEPTIDE
     332 <222> LOCATION: (2)
     333 <223> OTHER INFORMATION: where X is any aa
     335 <400> SEQUENCE S
W--> 336 Leu Xaa Xaa Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Xaa
         ~
     337
W--> 339/Xaa Xaa Xaa Xaa Xaa Leu (Xaa Xaa Xaa Xaa Xaa Xaa Xaa Leu
     340
                      20
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DATE: 07/12/2000

RAW SEQUENCE LISTING

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

DATE: 07/12/2000

PATENT APPLICATION: US/09/606,129

TIME: 10:58:15

Input Set : A:\U607921.app

Output Set: N:\CRF3\07122000\1606129.raw

L:11 M:270 C: Current Application Number differs, Replaced Application Number L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:284 M:341 W: (46) "n" or "Xaa" used. for SEQ ID#:6
L:303 M:341 W: (46) "n" or "Xaa" used. for SEQ ID#:7
L:336 M:341 W: (46) "n" or "Xaa" used. for SEQ ID#:9
L:339 M:341 W: (46) "n" or "Xaa" used. for SEQ ID#:9
L:386 M:341 W: (46) "n" or "Xaa" used. for SEQ ID#:12
L:433 M:341 W: (46) "n" or "Xaa" used. for SEQ ID#:15
L:452 M:341 W: (46) "n" or "Xaa" used. for SEQ ID#:16
L:471 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17